



SEQUENCE LISTING

<110> James, Kenneth D.
Rahdakrishnan, Balasingham
Malkar, Navdeep B.
Miller, Mark A.
Ekwuribe, Nnochiri N.

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<130> 9233.108

<140> US 10/723, 933
<141> 2003-11-26

<150> US 60/429, 151
<151> 2002-11-26

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<170> PatentIn version 3.2

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35 40 45

His Leu Gln Gly Lys Leu Ser Glu Leu Gln Val Glu Gln Thr Ser Leu
50 55 60

Glu Pro Leu Gln Glu Ser Pro Arg Pro Thr Gly Val Trp Lys Ser Arg
65 70 75 80

Glu Val Ala Thr Glu Gly Ile Arg Gly His Arg Lys Met Val Leu Tyr
85 90 95

Thr Leu Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser Gly Cys
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Lys Val Leu Arg Arg His
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20

25

30

Gly Ser Ala Ser Asp Leu Glu Thr Ser Gly Leu Gln Glu Gln Arg Asn
35 40 45

His Leu Gln Gly Lys Leu Ser Gu Leu Gln Val Glu Gln Thr Ser Leu
50 55 60

Glu Pro Leu Gln Glu Ser Pro Arg Pro Thr Gly Val Trp Lys Ser Arg
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Thr Leu Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser Gly
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<400> 48

Xaa Val Leu Arg
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<400> 49

Xaa Val Leu Arg Xaa
1 5

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<223> Xaa may be Arg or Lys

<220>
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<222> (6)..(6)
<223> Xaa may be Tyr or His

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Xaa Val Leu Arg Xaa Xaa
1 5

<210> 51
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<223> Xaa cannot be Asn if amino acid 25 is Arg and amino acid 26 is Tyr

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<222> (25)..(25)
<223> Xaa cannot be Arg if amino acid 21 is Asn and amino acid 26 is Tyr

<220>
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<222> (26)..(26)
<223> Xaa cannot be Tyr if amino acid 21 is Asn and amino acid 25 is Arg

<400> 51

Asp Ser Gly Cys Phe Gly Arg Arg Leu Asp Arg Ile Gly Ser Leu Ser
1 5 10 15

Gly Leu Gly Cys Xaa Val Leu Arg Xaa Xaa
20 25

<210> 52
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<400> 52

Asn Val Leu Arg Arg Tyr
1 5

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<223> Polypeptide may be present or absent

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<222> (7)..(9)
<223> Polypeptide may be present or absent

<400> 53

Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His
20 25 30

<210> 54
<211> 9
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<400> 54

Ser Pro Lys Met Val Gln Gly Ser Gly
1 5

<210> 55
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<220>
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<400> 55

His His His His His Ala Asp Gly Glu
1 5 10

<210> 56
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<220>
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<400> 56

Ala Asp Gly Glu
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<210> 57
<211> 8
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<220>
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<400> 57

Arg Arg Asp Ala Glu Asp Pro Arg
1 5

<210> 58
<211> 5
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<213> Artificial sequence

<220>
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<400> 58

Glu Gly Asp Arg Arg
1 5

<210> 59
<211> 11
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<220>
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<400> 59

His His His His His Glu Gly Asp Arg Arg
1 5 10

<210> 60
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<400> 60

Arg Arg Asp Ala Glu Asp Arg Arg
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<210> 61

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<223> Xaa can be any naturally occurring amino acid

<400> 61

His His His His His Xaa Glu Gly Asp Arg Arg
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<210> 62
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<220>
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<400> 62

Arg Gly Asp Ala Glu Asp Pro Arg
1 5

<210> 63
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<400> 63

Glu Gly Asp Pro Arg
1 5

<210> 64
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<400> 64

His His His His His Glu Gly Asp Pro Arg
1 5 10

<210> 65
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<400> 65

Ala Arg Gly Asp Ala Glu Asp Pro Arg
1 5

<210> 66
<211> 9
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<222> (7)..(7)
<223> Xaa can be any naturally occurring amino acid

<400> 66

His His His His His Xaa Met Met
1 5

<210> 67
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<400> 67

Asp Asp Ala Gly Glu
1 5

<210> 68
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<400> 68

His His His His His Ala Asp Gly Glu
1 5 10

<210> 69
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<400> 69

Glu Ala Gly Glu
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<210> 70
<211> 4
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<400> 70

Glu Gly Asp Ala
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<210> 71
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<400> 71

Glu Gly Asp Ala His His His His His Glu
1 5 10

<210> 72
<211> 11
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<400> 72

Glu His His His His His Ala Asp Gly Glu
1 5 10

<210> 73

<211> 32

<212> PRT

<213> Homo sapiens

<220>

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<222> (10)..(26)

<223> Disulfide bond may be present or absent

<400> 73

Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His
20 25 30

<210> 74

<211> 32

<212> PRT

<213> Homo sapiens

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<221> MOD_RES

<222> (1)..(1)

<223> A modifying moiety may be present

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<221> MISC_FEATURE

<222> (31)..(31)

<223> Xaa is not Arg

<400> 74

Thr Ala Pro Arg Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Met
1 5 10 15

Asp Arg Ile Gly Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Xaa Tyr
20 25 30

<210> 75

<211> 32

<212> PRT
<213> Canis familiaris

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<223> Xaa can be any naturally occurring amino acid

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<223> Xaa can be any naturally occurring amino acid

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<223> Xaa can be any naturally occurring amino acid

<400> 75

Ser Pro Xaa Met Met His Xaa Gly Gly Cys Phe Gly Arg Arg Leu Asp
1 5 10 15

Arg Ile Gly Ser Leu Ser Gly Leu Gly Cys Asn Val Leu Arg Xaa Tyr
20 25 30

<210> 76
<211> 38
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<213> Homo sapiens

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<223> Xaa can be any naturally occurring amino acid

<400> 76

Glu Val Xaa Tyr Asp Pro Cys Phe Gly His Xaa Ile Asp Arg Ile Asn
1 5 10 15

His Val Ser Asn Leu Gly Cys Pro Ser Leu Arg Asp Pro Arg Pro Asn
20 25 30

Ala Pro Ser Thr Ser Ala
35

<210> 77
<211> 22
<212> PRT
<213> Homo sapiens

<400> 77

Gly Leu Ser Lys Gly Cys Phe Gly Leu Lys Leu Asp Arg Ile Gly Ser
1 5 10 15

Met Ser Gly Leu Gly Cys
20

<210> 78
<211> 28
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<222> (12)..(12)
<223> Xaa can be any naturally occurring amino acid

<400> 78

Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Xaa Asp Arg Ile Gly
1 5 10 15

Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr
20 25

<210> 79
<211> 17
<212> PRT
<213> Artificial sequence

<220>
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<222> (5)..(5)
<223> Xaa may be any amino acid other than Lys

<400> 79

Cys Phe Gly Arg Xaa Met Asp Arg Ile Ser Ser Ser Gly Leu Gly
1 5 10 15

Cys

<210> 80
<211> 36
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<223> Xaa may be any naturally occurring amino acid, and may be present or absent

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<223> Xaa may be any naturally occurring amino acid, and may be present or absent

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<223> Xaa may be any naturally occurring amino acid, and may be present or absent

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<223> Xaa may be any naturally occurring amino acid, and may be present or absent

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<222> (35)..(35)
<223> Xaa can be any naturally occurring amino acid

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<222> (36)..(36)
<223> Xaa may be any naturally occurring amino acid, and may be present or absent

<400> 80

Ser Pro Arg Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Xaa Xaa Xaa Xaa Xaa Xaa
20 25 30

Xaa Xaa Xaa Xaa
35

<210> 81
<211> .6
<212> PRT
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<220>
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<400> 81

Arg Val Leu Arg Arg His
1 5

<210> 82
<211> 32
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<222> (14)..(14)
<223> Xaa may be any amino acid other than Lys

<220>
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<222> (27)..(27)
<223> Xaa can be any naturally occurring amino acid

<400> 82

Ser Pro Xaa Met Val Gln Gly Ser Gly Cys Phe Gly Arg Xaa Met Asp
1           5           10          15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Xaa Val Leu Arg Arg His
20           25          30

<210> 83
<211> 32
<212> PRT
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<222> (14)..(14)
<223> Xaa is not Lys

<220>
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<222> (27)..(27)
<223> Xaa is not Lys

<400> 83

Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Xaa Met Asp
1           5           10          15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Xaa Val Leu Arg Arg His
20           25          30

<210> 84
<211> 19
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<213> Artificial sequence
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<222> (19)..(19)
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<400> 84

Xaa Cys Phe Gly Arg Arg Met Asp Arg Ile Ser Ser Ser Ser Gly Leu
1 5 10 15

Gly Cys Xaa

<210> 85
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<212> PRT
<213> Artificial sequence

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<400> 85

Ser Pro Lys Met Val Gln Gly Ser Gly Cys
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<210> 86
<211> 9
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<400> 86

Pro Lys Met Val Gln Gly Ser Gly Cys
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<210> 87
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<400> 87

Lys Met Val Gln Gly Ser Gly Cys
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<210> 88
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<400> 88

Met Val Gln Gly Ser Gly Cys
1 5

<210> 89
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<400> 89

Val Gln Gly Ser Gly Cys
1 5

<210> 90
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<400> 90

Gln Gly Ser Gly Cys
1 5

<210> 91
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<400> 91

Gly Ser Gly Cys
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<210> 92
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Ser Pro Lys Met
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<210> 93
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Ser Pro Lys Met Val
1 5

<210> 94
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<400> 94

Ser Pro Lys Met Val Gln
1 5

<210> 95
<211> 4
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<400> 95

Lys Met Val Gln
1

<210> 96
<211> 5
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<400> 96

Lys Met Val Gln Gly
1 5

<210> 97
<211> 6
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<400> 97

Lys Met Val Gln Gly Ser
1 5

<210> 98
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<400> 98

Lys Met Val Gln Gly Ser Gly
1 5

<210> 99
<211> 8
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Lys Met Val Gln Gly Ser Gly Cys
1 5

<210> 100

<211> 6

<212> PRT

<213> Artificial sequence

<220>

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<400> 100

Lys Val Leu Arg Arg His
1 5

<210> 101

<211> 5

<212> PRT

<213> Artificial sequence

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<400> 101

Lys Val Leu Arg Arg
1 5

<210> 102

<211> 4

<212> PRT

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Lys Val Leu Arg
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<210> 103

<211> 6

<212> PRT

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<400> 103

Arg Val Leu Arg Arg His
1 5

<210> 104

<211> 5

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Arg Val Leu Arg Arg
1 5

<210> 105

<211> 4

<212> PRT

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<400> 105

Arg Val Leu Arg
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<210> 106

<211> 29

<212> PRT

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<223> Xaa is not Lys

<400> 106

Ser Pro Xaa Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Lys Val Leu
20 25

<210> 107
<211> 26
<212> PRT
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<223> Xaa is not Lys

<400> 107

Ser Pro Xaa Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Ser Gly Leu Gly Cys
20 25

<210> 108
<211> 33
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<223> Xaa may be any naturally occurring amino acid and may be present or absent

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<223> Xaa may be any naturally occurring amino acid and may be present

or absent

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<223> Xaa may be any naturally occurring amino acid and may be present or absent

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<223> Xaa may be any naturally occurring amino acid and may be present or absent

<400> 108

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Phe Gly Arg Arg Met
1 5 10 15

Asp Arg Ile Ser Ser Ser Gly Leu Gly Cys Arg Val Leu Arg Arg
20 25 30

His

<210> 109
<211> 17
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<220>
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<222> (5)..(5)
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<220>
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<222> (10)..(10)
<223> Xaa may be Ser or Lys

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa is Ser and may be present or absent

<220>
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<222> (12)..(12)
<223> Xaa is Ser and may be present or absent

<220>
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<222> (13)..(13)
<223> Xaa is Ser and may be present or absent

<400> 109

Cys Phe Gly Arg Xaa Met Asp Arg Ile Xaa Xaa Xaa Xaa Gly Leu Gly
1 5 10 15

Cys

<210> 110
<211> 32
<212> PRT
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<220>
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<220>
<221> MISC_FEATURE
<222> (30)..(30)
<223> Xaa is not Arg

<400> 110

Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Lys Val Arg Xaa Arg His
20 25 30

<210> 111
<211> 32
<212> PRT
<213> Artificial sequence

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<222> (27)..(27)
<223> Xaa is not Lys

<400> 111

Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Xaa Val Leu Arg Arg His
20 25 30

<210> 112
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<222> (33)..(33)
<223> Xaa may be Lys or Cys

<400> 112

Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His
20 25 30

Xaa

<210> 113
<211> 26
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<220>
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<223> Xaa is not Lys

<220>
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<223> Xaa may be Gly, Met, Leu, Phe, Ile, or a conservative substitution thereof

<220>
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<222> (24)..(24)
<223> Xaa may be Leu, Trp, Tyr, Phe, or a conservative substitution thereof

<220>
<221> MISC_FEATURE
<222> (25)..(25)
<223> Xaa may be Gly, Arg, or a conservative substitution thereof

<400> 113

Ser Pro Xaa Met Val Gln Gly Ser Gly Cys Phe Gly Arg Xaa Met Asp
1 5 10 15

Arg Ile Ser Ser Ser Xaa Xaa Xaa Cys
20 25

<210> 114
<211> 23
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<213> Artificial sequence

<220>
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<221> MISC_FEATURE
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<223> Xaa may be Thr, Ala, Arg, His, Pro, or Glu

<220>
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<222> (12)..(12)
<223> Xaa may be Lys, Asn, Arg, Ser, Asp, or Pro

<220>
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<222> (12)..(12)
<223> Methylation if Xaa is Asn

<220>
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<222> (17)..(17)
<223> Xaa is not Gly

<220>
<221> MOD_RES
<222> (17)..(17)
<223> Xaa may be Orn, Har, p-amidinophenyl Ala, or Ile

<400> 114

Lys Cys Phe Lys Gly Lys Asn Asp Arg Xaa Lys Xaa Gln Ser Gly Leu
1 5 10 15

Xaa Cys Asn Ser Phe Lys Tyr
20

<210> 115
<211> 195
<212> PRT
<213> Artificial sequence

<220>
<223> BNP pro-pentapeptide

<400> 115

His His His His His Glu Gly Asp Arg Arg Ser Pro Lys Met Val
1 5 10 15

Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp Arg Ile Ser Ser Ser
20 25 30

Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His Arg Asp Ala Glu
35 40 45

Asp Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met

50

55

60

Asp Arg Ile Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg
65 70 75 80

His Arg Arg Asp Ala Glu Asp Ser Pro Lys Met Val Gln Gly Ser Gly
85 90 95

Cys Phe Gly Arg Lys Met Asp Arg Ile Ser Ser Ser Gly Leu Gly
100 105 110

Cys Lys Val Leu Arg Arg His Arg Arg Asp Ala Glu Ap Ser Pro Lys
115 120 125

Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp Arg Ile Ser
130 135 140

Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His Arg Arg Asp
145 150 155 160

Ala Glu Asp Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg
165 170 175

Lys Met Asp Arg Ile Ser Ser Ser Gly Leu Gly Cys Lys Val Leu
180 185 190

Arg Arg His
195

<210> 116

<211> 29

<212> PRT

<213> Artificial sequence

<220>

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Xaa Xaa Xaa Xaa Xaa
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Gln Gly Ser Gly
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Cys Lys

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Arg Ile Ser Ser Ser Ser Gly Leu Gly Cys Xaa Val Leu Arg Arg His
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Xaa Xaa Xaa Xaa Xaa
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